

Art
Art
b) generating recombinant libraries comprising said RRT-SH3 domains, and

c) subjecting said libraries to affinity or functional selection steps to identify artificial SH3 domains.

Subst
02
17. (amended) The method according to claim 3, wherein the six amino acids that are replaced in the RT-loop are replaced with a peptide motif derived from Hck-SH3 and targeted to the HIV-1 Nef protein selected from the group consisting of XSWSXX (SEQ ID NO:28), XSPFXX (SEQ ID NO:30) and XSXFPW (SEQ ID NO:32), wherein X is any amino acid.
